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Review

Necroptosis: The Trojan horse in cell autonomous antiviral host defense



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ARTICLE INFO

Article history:

Received 4 February 2015
Returned to author for revisions
12 February 2015
Accepted 2 March 2015
Available online 24 March 2015

Keywords:

Programmed necrosis
FADD
cFLIP
Serine/threonine protein kinase
Ribonucleotide reductase

ABSTRACT

Herpesviruses suppress cell death to assure sustained infection in their natural hosts. Murine cytomegalovirus (MCMV) encodes suppressors of apoptosis as well as M45-encoded viral inhibitor of RIP activation (vIRA) to block RIP homotypic interaction motif (RHIM)-signaling and recruitment of RIP3 (also called RIPK3), to prevent necroptosis. MCMV and human cytomegalovirus encode a viral inhibitor of caspase (Casp)8 activation to block apoptosis, an activity that unleashes necroptosis. Herpes simplex virus (HSV)1 and HSV2 incorporate both RHIM and Casp8 suppression strategies within UL39-encoded ICP6 and ICP10, respectively, which are herpesvirus-conserved homologs of MCMV M45. Both HSV proteins sensitize human cells to necroptosis by blocking Casp8 activity while preventing RHIM-dependent RIP3 activation and death. In mouse cells, HSV1 ICP6 interacts with RIP3 and, surprisingly, drives necroptosis. Thus, herpesviruses have illuminated the contribution of necroptosis to host defense in the natural host as well as its potential to restrict cross-species infections in nonnatural hosts.

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Introduction

Necroptosis is a form of programmed cell death leading to membrane leakage independent of caspases, orchestrated by the activity of receptor interacting protein (RIP)3 kinase and its target, a pseudokinase called mixed lineage kinase-like (MLKL). Hallmark insights using vaccinia virus raised the specter of RIP3 kinase-dependent necroptosis contributing to antiviral host defense (Cho et al., 2009). The identification of a specific viral inhibitor of RIP

activation (vIRA) encoded by the herpesvirus murine cytomegalovirus (MCMV) demonstrated that RIP3-mediated necroptosis is a *bona fide* cell autonomous host defense pathway subjected to a specific viral countermeasure (Upton et al., 2010, 2012). Necroptosis has been implicated in additional viral (Berger and Danthi, 2013; Guo et al., 2015) as well as bacterial (Robinson et al., 2012; Weng et al., 2014) infections and is probably a default pathway when caspase (Casp)8 activity is compromised in cells with sufficient levels of RIP3 (Upton and Chan, 2014; Kaiser et al., 2013). Many viruses, including both poxviruses and herpesviruses, inhibit caspase (Casp)8 activity (Mocarski et al., 2011) and therefore have the potential to unleash this alternate death pathway. For example, the betaherpesvirus-conserved viral inhibitor of Casp8 activation (vICA) naturally suppresses virus-induced apoptosis in macrophages (Menard et al.,

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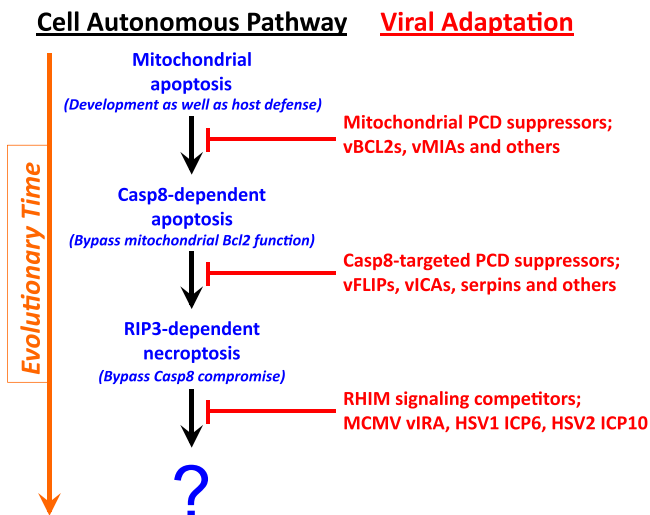


Fig. 1. Evolutionary relationships in cell autonomous death pathways and virus-encoded countermeasures. Programmed cell death (PCD) pathways include the mitochondrial (or cell-intrinsic) death pathway that eliminates cells during development and homeostatic turnover as well as in host defense (Kaiser et al., 2013; Mocarski et al., 2011; McCormick and Mocarski, 2013; Galluzzi et al., 2008). Viral (v)Bcl2 homologs and other virus-encoded mitochondrial suppressors block mitochondrial PCD, establishing conditions that selected for a Casp8-like activity to directly trigger effector caspases, bypassing mitochondrial Bcl2 function. Viruses adaptation included Casp8-targeted cell death suppressors, some specific, like herpesvirus vFLIPs, betaherpesvirus vICA, and some nonspecific, like the poxvirus serpin, CrmA. RIP3-dependent necroptosis via RHIM-dependent interaction with RIP1, TRIF or DAI, enables the host to bypass the impact of virus-encoded Casp8 inhibitors. The examples of MCMV vIRA and HSV1/HSV2 RHIM signaling competitors are the first examples of viral suppressors of necroptosis (Guo et al., 2015; Upton et al., 2008, 2010, 2012). Note that HSV1 ICP6 inhibits necroptosis in human cells but promotes necroptosis in mouse cells (Huang et al., 2015; Wang et al., 2014). This depiction is adapted from (Kaiser et al., 2013).

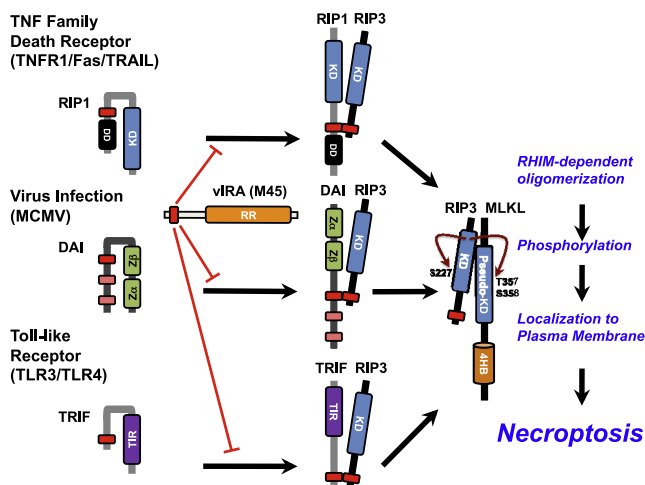


Fig. 2. Canonical and noncanonical RHIM-dependent activation of RIP3 kinase. Canonical activation of necroptosis via RIP1 recruitment and activation RIP3 downstream of TNF family death receptors (TNFR1, Fas, TRAIL), highlighting the known and predicted conformational changes that accompany this pathway (Chan et al., 2014). Noncanonical activation of necroptosis via DAI downstream of MCMV infection or via TRIF downstream of TLR3 and TLR4 activation, driving recruitment and activation of RIP3, with predicted conformational changes. All forms of RHIM-dependent activation of RIP3 are blocked by MCMV-encoded vIRA and converge on the RIP3-dependent recruitment and phosphorylation events leading to plasma membrane leakage and other outcomes characteristic of necroptosis.

2003; McCormick et al., 2003, 2010; Skaletskaya et al., 2001; Cicin-Sain et al., 2008). MCMV relies on M45-encoded vIRA to prevent necroptosis and sustain infection. The M45 protein is an enzymatically inactive homolog of the large subunit (R1) of ribonucleotide

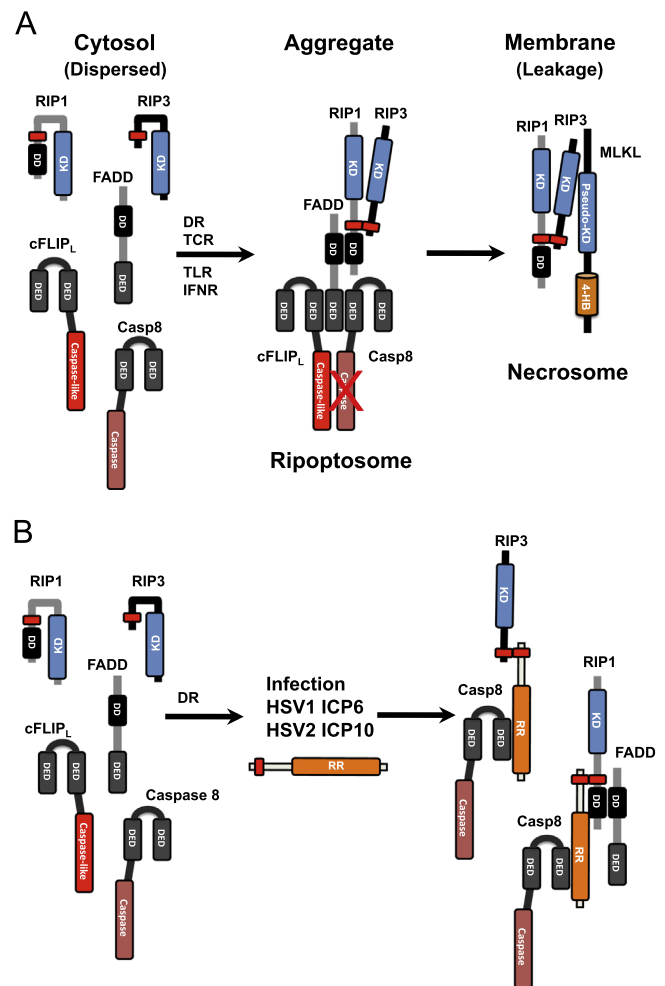


Fig. 3. Formation and disruption of the ripoptosome and necrosome by HSV1 ICP6 and HSV2 ICP10. (A) The ripoptosome forms from cytosolic components. FADD becomes activated downstream of TNF family death receptor (DR) activation, as well as following T cell receptor (TCR), TLR and interferon receptor (IFNR) activation, to recruit Casp8 and cFLIP_L heterodimer via homotypic death effector domain (DED) as well as RIP1 via a homotypic death domain (DD) interactions. RIP1 recruits RIP3 via a common RHIM (shown as a red rectangle) and, when Casp8 activity is inhibited, triggers oligomerization that leads to an amyloid-like complex that recruits MLKL into a necrosome that localizes to membranes and directs the final steps in necroptosis leading to membrane leakage. (B) Interaction of HSV1 ICP6 and HSV2 ICP10 with ripoptosome components in human cells. Cytosolic ripoptosome components form HSV R1-dependent complexes via RHIM and Casp8 interaction domains discussed in the text.

reductase (RNR) common across the herpesviruses. Based on studies of MCMV, we have posited that an evolutionary dialog has gone on between host cell death pathways and virus-encoded cell death suppressors that is evident in mammals (Kaiser et al., 2013; Mocarski et al., 2014), as depicted in Fig. 1. Mitochondrial, or cell-intrinsic cell death is an ancient form of host defense as well as a common target of virus-encoded suppressors that provide the selective pressure for Casp8 extrinsic apoptosis to side-step mitochondria and directly activate the executioner caspases, Casp3 and Casp7. The ability of a self-activating caspase to execute cells prompted the acquisition by viruses of specific Casp8 inhibitors (Mocarski et al., 2011). This selected for host adaptation of necroptosis as a trap door to eliminate viruses that suppress Casp8, built upon established RIP homotypic interaction motif (RHIM) signaling pathways. RHIM-competitors encoded by MCMV, HSV1 and HSV2 thus represent most recent evolutionary adaptation in this ancient pathogen-host stand-off (Kaiser et al., 2013; Mocarski et al., 2014).

MCMV vIRA employs an amino-terminal RHIM to interrupt signal transduction leading to cell death and cytokine activation. In

mice or humans, RIP1, DNA-induced activator of interferon (DAI) and TIR-domain-containing adapter-inducing interferon β (TRIF) can partner with RIP3, as depicted in Fig. 2 (Upton et al., 2010, 2012; Rebsamen et al., 2009; He et al., 2011; Kaiser et al., 2013). The MCMV-encoded RHIM competitor, vIRA prevents activation of RIP3 kinase activity, the subsequent recruitment of MLKL, and the later steps leading to cell leakage (Kaiser et al., 2013). vIRA blocks all consequences of RHIM-dependent signaling, including activation of NF- κ B (Rebsamen et al., 2009; Upton et al., 2008; Kaiser et al., 2008) even though the impact on this transcription factor does not appear to impact infection in mouse cells or mice. Despite the crucial role that MCMV M45-encoded vIRA plays in cell death suppression, and the likely preservation of this mechanism in other rodent betaherpesviruses, the HCMV UL45 is inactive (Hahn et al., 2002; Patrone et al., 2003). Nevertheless, HCMV blocks necroptosis during infection by employing a strategy that subverts a later step in the pathway (Omoto et al., 2015). The M45 homologs of herpes simplex virus (HSV)1 and HSV2 (UL39 gene products, ICP6 and ICP10, respectively) have recently been shown to employ a RHIM competitor mechanism in human cells (Guo et al., 2015). Although analogous to MCMV vIRA, HSV cell death suppressors ICP6 and ICP10 make use of an anti-apoptotic domain separate from the RHIM to interact directly with Casp8 death effector domains (Dufour et al., 2011), a function mediated by the carboxyl-terminal RNR domain (Guo et al., 2015). Here, we will focus on examples of virus-encoded RHIM-signaling inhibitors and the contribution of Casp8 inhibition to unleashed necroptosis.

Activation and suppression of necroptosis

Necroptosis hinges on RIP3 RHIM-mediated signal transduction that was initially characterized in cells undergoing cell death following stimulation with TNF in the presence of caspase 8 inhibitor (Fig. 3A) or in cells that lack Casp8 (Cho et al., 2009; He et al., 2009; Zhang et al., 2009), and is the subject of up-to-date reviews (Sun and Wang, 2014; Chan et al., 2014). RIP3 is a serine-threonine kinase that recruits and phosphorylates the pseudokinase MLKL, controlling oligomerization events at membranes that lead to cellular changes including leakage of cytosolic contents (Sun et al., 2012; Zhao et al., 2012; Murphy et al., 2013; Chen et al., 2013; Hildebrand et al., 2014). RHIM-dependent interaction of RIP1 and RIP3 characterizes the canonical form of this death pathway (Cho et al., 2009; He et al., 2009; Zhang et al., 2009) and results in conformational changes and amyloid-like oligomerization of a RIP1–RIP3 complex (Chan et al., 2014; Li et al., 2012). Canonical TNFR1-mediated necroptosis requires the protein kinase activity of RIP1 in addition to RHIM interaction with RIP3. RIP3 activation and execution of necroptosis may proceed independent of RIP1. DAI as well as TRIF can naturally partner with RIP3 and trigger a noncanonical necroptosis, independent of RIP1 RHIM and kinase activities (Upton et al., 2010, 2012; Kaiser et al., 2013). Interestingly, experiments have substantiated the noncanonical pathway even downstream of death receptors (Fig. 1) (Moujalied et al., 2013; McQuade et al., 2013; Cho et al., 2011). Regardless of the involvement of RIP1, RIP3 must autophosphorylate itself to facilitate interaction with and subsequent phosphorylation of MLKL (Kaiser et al., 2013; Chan et al., 2014; Wu et al., 2013) (Fig. 2). A cytosolic Casp8-containing complex, called Complex IIB (Micheau and Tschopp, 2003) or the ripoptosome (Chan et al., 2014; Tenev et al., 2011; Feoktistova et al., 2012, 2011), forms downstream of TNFR1 or of TRIF-dependent signaling pathways controlled by TLR3 or TLR4. This complex is increasingly recognized to control apoptotic as well as necroptotic outcomes (Chan et al., 2014; Mandal et al., 2014) after assembling from preexisting cytosolic components in a manner that is independent of pro-apoptotic caspase activity as well as mitochondrial function (Tait et al., 2013). The ripoptosome is assembled around Casp8–cFLIP_L heterodimer in association with FADD to which

RIP1 and RIP3 are recruited through well-established protein–protein homotypic interaction domains depicted in Fig. 3A. This assembly has been the subject of many reviews (Sun and Wang, 2014; Chan et al., 2014; van Raam and Salvesen, 2012). MLKL is recruited and phosphorylated by RIP3 kinase within a necrosome complex (Murphy et al., 2013; Hildebrand et al., 2014; Cai et al., 2014; Wang et al., 2014; Dondelinger et al., 2014) resulting in MLKL conformational changes and oligomerization that result in membrane association, as well as the pattern of disruption and leakage (Sun and Wang, 2014; Chan et al., 2014).

Casp8 inhibition has long been recognized as a key factor in the induction of necrotic death (Holler et al., 2000; Vercammen et al., 1998) and is a crucial feature in terms of how this pathway is unleashed in virus-infected cells (Kaiser et al., 2013; Mocarski et al., 2011). Besides suppression of apoptosis, inhibition of Casp8 also fundamentally affects the availability of key ripoptosome components. The susceptibility of RIP1 (Lin et al., 1999) and RIP3 (Feng et al., 2007) to Casp8 cleavage-dependent inactivation of pro-necrotic kinase activity suggests that both of these pro-necrotic kinases are silenced by Casp8 (Sun and Wang, 2014; Chan et al., 2014). Although the heterodimeric complex of Casp8 and cFLIP_L elaborates a basal protease activity that may be sufficient to inactivate RIP1 and/or RIP3, details of this regulatory process have not yet been worked out. Whereas Casp8 suppression by vICA is needed during natural infection to block apoptosis in macrophages (Cicin-Sain et al., 2008), MCMV-infected cells become sensitized to necroptosis such that vIRA is necessary to suppress the RHIM-dependent recruitment of RIP3 by DAI in all infected host tissues (Upton et al., 2010, 2012). Without an intact RHIM in vIRA, MCMV cannot sustain infection in wild-type mice. However, normal infection is restored in RIP3- or DAI-deficient strains of mice. Mice carrying kinase inactive RIP3 also normalize replication of vIRA-deficient MCMV (Mandal et al., 2014), consistent with the essential contribution of this pro-necrotic kinase activity to host defense. The investigation of cell death suppressor mutants of MCMV has revealed the importance of deflecting both Casp8-mediated apoptosis and RIP3-mediated necroptosis (McCormick and Mocarski, 2013). Necroptosis is triggered in tissue-resident cells regardless of whether the same cells are sensitive to apoptosis. Importantly, studies with vIRA-deficient MCMV have revealed that necroptosis may be unleashed in any virus-infected cell; however, mouse or human cells placed into culture vary in their levels of RIP3 and therefore vary in their susceptibility to this death pathway. The reasons why cultured cells tend to lose RIP3 (as well as MLKL) expression is presently unknown, but certainly suggests that cell culture selects against the retention of necroptotic machinery. In addition to herpesviruses, this alternate death is likely to be modulated by poxviruses and other DNA viruses known to block Casp8 activity (Mocarski et al., 2011; Hedrick et al., 2010).

Cell death outcomes are strongly influenced by posttranslational modification that acts as a scaffold such as those provided by K63 or M1 (linear) polyubiquitination (Silke and Meier, 2013; Walczak, 2011). RIP1 seems to be the most important target of such modifications, in line with its contribution to balancing death outcomes emanating from the ripoptosome (Feoktistova et al., 2012; Walczak, 2011; Kaiser et al., 2014; Weinlich and Green, 2014). Compromised K63 or linear polyubiquitination increases susceptibility of cells to apoptotic as well as necroptotic death (Kaiser et al., 2013). Inhibitors and mutants affecting either of two distinct E3 ubiquitin ligase complexes, cellular inhibitor of apoptosis (cIAP)1/cIAP2 (Silke and Meier, 2013) or linear ubiquitination chain assembly complex (LUBAC) (Walczak, 2011) have been employed to show the importance of polyubiquitination in dictating sensitivity to death signaling (He et al., 2009; Tenev et al., 2011; Feoktistova et al., 2011; Vanlangenakker et al., 2011; Varfolomeev et al., 2007; Gerlach et al., 2011). Although the polyubiquitination machinery is not a known target of mammalian virus-encoded cell death

suppressors, the first IAP ever identified emerged from studies in baculoviruses where virus-encoded IAPs suppress cell death to sustain infection (Crook et al., 1993; Birnbaum et al., 1994). To date only large DNA viruses infecting arthropods encode IAP-like suppressors of apoptosis (Clem, 2005). Interestingly, the noncanonical induction of necroptosis that occurs in MCMV M45 mutant virus infected cells is insensitive to IAP1/IAP2 antagonists, consistent with known RIP1 independent characteristics of the pathway (Upton et al., 2010, 2012). Only time and additional investigation will reveal whether mammalian and insect cells differ in some fundamental way in harnessing IAP function for host defense.

Uncontrolled necroptosis undermines development and tissue homeostasis

The discovery that midgestational death in Casp8 or FADD-deficient mice results from unrestricted RIP3- and RIP1-dependent death (Oberst et al., 2011; Kaiser et al., 2011; Zhang et al., 2011) completely revised the thinking about ripoptosome function in development and inflammatory disease. These investigations, together with incisive observations on Casp8- or FADD-deficient T cells (Lu et al., 2011; Ch'en et al., 2011) revealed how dysregulation of death pathways during development as well as during the T cell response to antigen impact cell fate. The complex dialog between FADD/Casp8 and pro-necrotic kinases is further bolstered by an investigation of FADD-deficient keratinocytes (Bonnet et al., 2011) and intestinal epithelium (Welz et al., 2011) as well as in a number of other settings (Mocarski et al., 2011, 2014; Sun and Wang, 2014; Chan et al., 2014; Weinlich et al., 2011). Thus, it is now widely understood that midgestational death due to germ line deficiency in Casp8 (or FADD) is due to canonical RIP1–RIP3 necroptosis, particularly because the phenotype is reversed by either kinase-inactive RIP3 (Mandal et al., 2014; Newton et al., 2014) or kinase-inactive RIP1 (Kaiser et al., 2014). Both RIP1 and RIP3 kinase activities contribute directly to fetal demise. Consistent with the observations on Casp8 and FADD, serious defects are observed with germ line deficiency in other ripoptosome components. cFLIP-deficient (Dillon et al., 2012) or RIP1-deficient mice (Kaiser et al., 2014; Dillon et al., 2014; Rickard et al., 2014) die during development but are fully rescued by the combined elimination of apoptotic (Casp8 or FADD) and necroptotic (RIP3 or MLKL) machinery. Thus, mice deficient in multiple complementary components are fully viable, fertile, and where examined (Kaiser et al., 2014), immunocompetent and able to clear virus infection. The picture that emerges from these studies and from a growing number of examples where disruption of Casp8 is performed in a tissue-specific manner (Sun and Wang, 2014; Chan et al., 2014) is that the balance of Casp8-dependent apoptosis and RIP3-dependent necroptosis is crucial to sustain tissue homeostasis (Mocarski et al., 2014; Chan et al., 2014). The common ability of death receptors, TLRs, T cell receptors, genotoxic stress and interferon receptors as well as virus infection to drive innate immune cell death as well as innate immune cytokine activation indicates that the ripoptosome may act as a pathogen supersensor to guard mammalian cells from insult (Kaiser et al., 2013). Perturbation of the ripoptosome undermines development even though its function is dispensable for either development or tissue homeostasis.

It is important to note that developmental consequences of mutant ripoptosome components only show modest dependence on TNF signaling despite the fact that canonical, RIP1-dependent activation of RIP3 kinase activity and RHIM signal transduction are crucial to the demise of mutant mice (Cho et al., 2009; Kaiser et al., 2013, 2011; He et al., 2009; Zhang et al., 2009; Oberst et al., 2011). The ripoptosome forms under a wide variety of innate signaling conditions, executing either apoptosis or necroptosis (Kaiser et al., 2013). The recent demonstration that the ripoptosome is also triggered by conformational changes in RIP3 independent of its pro-necrotic kinase

activity (Mandal et al., 2014) reveals a remarkable capacity of this protein kinase to regulate caspase activation in addition to necroptosis. RIP3 deficiency itself does not cause any developmental problems (Newton et al., 2004) but reveals modest innate immune response dysregulation (Moriwaki et al., 2015). The potential for RIP3 kinase domain mutants as well as small molecule kinase inhibitors to trigger apoptosis dependent on RIP1 and RIP3 scaffold functions independent of activity of these kinases (Mandal et al., 2014; Newton et al., 2014) expanded the settings where the ripoptosome controls cell fate. Further dissection of the scaffold versus the enzymatic roles of RIP1 as well as RIP3 will provide further insights into the mechanism(s) underlying alternate death outcomes.

Evolutionary adaptation of innate immune programmed cell death in host defense

Evolutionary relationships of Casp8-mediated apoptosis and RIP3-mediated necroptosis pathways in response to viral suppressors have been reviewed (Kaiser et al., 2013) and are illustrated in Fig. 1. Our work with MCMV brought to light the importance of RIP3-dependent necroptosis as a host defense mechanism (Upton et al., 2012). Many diverse innate and adaptive immune mechanisms ranging from interferons to antibodies and T cells mediate control over pathogens, but only necroptosis has been shown to completely prevent a herpesvirus such as MCMV from infecting its natural host, even when high doses of virus are used. This can be attributed to the unique ability of necroptosis to eliminate virus-infected cells before production of progeny can occur, thereby shutting down the infection. Because the MCMV vIRA completely blocks the elaboration of necroptosis during MCMV infection, this remarkably potent host defense pathway becomes effectively neutralized. This absolutely fascinating pathogen–host relationship extends to human cell settings where necroptosis is targeted by HCMV (Omoto et al., 2015) as well as HSV gene products (Guo et al., 2015). In all of these settings, we hypothesize that Casp8, together with its partners, FADD and cFLIP_L, forms a pathogen supersensor (Kaiser et al., 2013) and triggers cell autonomous defense pathways that include cell death and cytokine production (Mocarski et al., 2014), although, in mice, death pathways predominate innate signaling and play out completely independent of interferon and inflammatory cytokine function. The subversion tactics by herpesviruses have codified the importance of neutralizing these pathways in pathogenesis.

Suppression of necroptosis by HSV

HSV1 suppresses programmed cell death induced by a wide variety of insults (Galvan and Roizman, 1998), relying on the regulatory protein ICP4 as well as protein kinase US3 (Leopardi and Roizman, 1996; Leopardi et al., 1997). In addition to US3 (Benetti and Roizman, 2004; Munger and Roizman, 2001), early viral gene products gD (Zhou et al., 2003; Zhou and Roizman, 2002) and, in particular, R1 (Dufour et al., 2011; Perkins et al., 2002a, 2002b; Langelier et al., 2002; Chabaud et al., 2007) are able to inhibit apoptosis, albeit in a cell-type specific manner (Han et al., 2009; Aurelian, 2005). HSV1 also manipulates apoptosis as a component of latency in rodent neurons (Perng et al., 2000; Du et al., 2012). The HSV1 and HSV2 R1 proteins, ICP6 and ICP10, suppress death receptor-dependent apoptosis by interacting with death effector domains of Casp8 via a conserved C-terminal RNR domain (Dufour et al., 2011). MCMV vIRA known to block apoptosis as well as necroptosis (Upton et al., 2008, 2010, 2012), but does not bind to Casp8 (Guo et al., 2015). HSV R1-mediated inhibition of Casp8 function may be analogous to betaherpesvirus vICA (McCormick et al., 2003; Skaletskaya et al., 2001). Until recently, little attention was given to necroptosis, although there have been

longstanding clues that caspase-independent pathways may dominate under certain conditions in HSV-infected cells and tissues (Aurelian, 2005; Galvan et al., 1999; Peri et al., 2008, 2011). Intriguingly, ICP6 and ICP10 exhibit sequence homology with the N-terminal RHIM of MCMV M45 and cellular RHIM-containing proteins (Lembo and Brune, 2009). This RHIM-like amino acid sequence is located in the amino-terminal extension of all R1 homologs encoded by the primate alphaherpesviruses and rodent betaherpesviruses. Recent comprehensive investigation of HSV1 ICP6 reveals an ability of this RHIM-like region to interact with RIP1 and RIP3 in human cells (Guo et al., 2015) as well as in mouse cells (Huang et al., 2015; Wang et al., 2014) (Fig. 3B); however, the RHIM-dependent interaction yields apparently opposite biological consequences in human versus mouse cells. Both HSV1 ICP6 and HSV2 ICP10 act as RHIM competitors in necroptosis-sensitive human cells (Guo et al., 2015), behavior that is reminiscent of MCMV vIRA (Upton et al., 2010, 2012). This behavior is consistent with a role in counteracting host defense. In contrast to the behavior in human cells, HSV1 ICP6 promotes necroptosis in mouse cells, where the RHIM-like region activates the pro-necrotic kinases (Huang et al., 2015; Wang et al., 2014). This is most dramatically demonstrated by comparing infection of parental WT HSV1 with an ICP6 mutant (Δ ICP6) HSV1. Infection with Δ ICP6 HSV1 mutant renders human cells highly sensitive to TNF due to the induction of apoptosis as well as necroptosis (Guo et al., 2015); whereas, infection with this mutant virus delays virus-induced necroptosis in mouse cells (Huang et al., 2015; Wang et al., 2014) (Fig. 3B). These findings raise important questions about the role of death pathways in modulating cross-species infections as well as about the reliance on rodents as surrogate hosts in studies of HSV1 pathogenesis and latency. It remains to be determined whether HSV2 ICP10 acts similarly to HSV1 ICP6 in mice. Importantly, and in contrast to the species-specific behavior of HSV1 ICP6, MCMV vIRA blocks either death receptor or virus-induced necroptosis function in human or mouse cells (Guo et al., 2015). Indeed, a swap of the M45 RHIM into HSV1 ICP6 converted the latter from an inducer to a suppressor of necroptosis in mouse cells (Huang et al., 2015).

The ability of HSV1 ICP6 and HSV2 ICP10 to bind Casp8 is integral to their suppression activity against necroptosis in human cells (Guo et al., 2015). Early (Dufour et al., 2011) and recent studies (Guo et al., 2015) demonstrate that Casp8 binding is mediated by the large carboxyl-terminal region of HSV R1 proteins, a region that is also responsible for ribonucleotide reductase activity. The amino-terminal region carrying the RHIM-like element is sufficient for interaction with RIP1 or RIP3 in a RHIM-dependent fashion, but the ability to block necroptosis in infected cells requires interaction with both the RIP1/RIP3 and Casp8-binding regions. Importantly, HSV1-infected cells are effectively sensitized to TNF-induced necroptosis by the Casp8-binding domain of ICP6 or ICP10. Importantly, ICP6mutRHIM virus-infected cells die prematurely following TNF treatment due to the suppression of Casp8 activity in the absence of RHIM competitor activity (Guo et al., 2015). Curiously, mouse cells are rescued from ICP6-induced necroptosis when infected with this same virus, confirming the contribution of RHIM signaling to the induction of the pathway in a species-specific fashion (Huang et al., 2015). From these recent studies, HSV1 ICP6 and HSV2 ICP10 emerge as important suppressors of necroptosis in cells from their natural host species, where Casp8 suppression sensitizes to necroptosis. This prevents RHIM signal competition, activation of RIP3 and recruitment of MLKL into a necrosome complex.

Perspectives

Pathogenesis studies in mice have unambiguously identified the cognate pathways targeted by virus-encoded cell death suppressors in natural pathogen-host settings, such as infection with MCMV, underscoring the importance of specific host defense mechanisms, including programmed cell death. Using MCMV mutants M45mutRHIM, both

RIP3 and DAI were found to be the cognate host cell targets of RHIM competitor activity. When they associate, necroptosis is activated to eliminate infected cells in all exposed tissues (Upton et al., 2008, 2010, 2012). These studies opened the way to show that RIP3 necroptosis underlies the phenotype of germ line Casp8-deficient mice (Kaiser et al., 2011) and that perinatal lethality due to combined effects of Casp8-dependent apoptosis and RIP3-dependent necroptosis underlies RIP1-deficiency (Kaiser et al., 2014). Along the way, we showed that the combined elimination of Casp8 and RIP3 programmed cell death pathways are remarkably dispensable for mammalian development, viability, fertility (Kaiser et al., 2014) and the ability to mount a fully developed T cell response to control viral infection (Kaiser et al., 2014). The key components of the ripoptosome become dispensable for development when eliminated in precise combinations that respect the ability of single mutants to dysregulate cell death by unleashing alternate cell death pathways, pathways that are elaborated in mammals to control pathogens. Recent demonstration that TNF-dependent necroptosis plays out against HSV1 in human cells and is blocked by the HSV1 and HSV2 R1 proteins adds significantly to the growing understanding of the antiviral role of necroptosis. These observations reveal viral evasion strategies that have the potential to prevent cross-species infections. The specter of necroptosis as a cross-species restriction factor, however, must be pursued further. Necroptosis emerges as a way to eliminate viruses capable of inhibiting Casp8, as illustrated initially by the behavior of vaccinia as well as by a growing body of data on MCMV infection and HSV1 infection, strategies that are likely to be usurped by other mammalian viruses.

Acknowledgments

Work performed in our laboratories is supported by NIH Grants to ESM (R01 AI020211) and WJK (DP1 OD012198), although the content is solely the responsibility of the authors and not the NIH or PHS.

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